## SEQUENCE LISTING

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•	_	cto	ctg	tee	_	tσσ	tac	200	acc		cta	acc	gac	cct		tet	96
		-	Leu		-		_				_						30
	Dea	Doc	Dou	20	DUI	.12	0,0	, II B	25	Oly	Dea	1114	пор	30		501	
	ctt	tgo	tat	gac	atc	acc	gtc	atc	cct	aag	ttc	aga	cct	gga	cca	cgg	144
	Leu	Cys	Tyr 35	Asp	Ile	Thr	Val	Ile 40	Pro	Lys	Phe	Arg	Pro 45	Gly	Pro	Arg	•
	tgg	tgt	gcg	gtt	caa	ggc	cag	gtg	gat	gaa	aag	act	ttt	ctt	cac	tat	192
			Ala														
		50					55					60					242
	-	-	ggc	-	-												240
	-	Cys	Gly	Ser	Lys		Val	Ihr	Pro	vai		Pro	Leu	Gly	Lys	Lys 80	
	65 cta	aat	gtc	aca	aca	70	taa	222	aca	cag	75 aac	cca	orta	cta	aga		288
			Val														200
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	vai	val	Asp	11e 100	Leu	Inr	GIU	GIN	105	Leu	Asp	116	GIN	110	Glu	Asn	
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	Tyr	Ilε	Pro 115	Lys	Glu	Pro	Leu	Thr 120	Leu	Gln	Ala	Arg	Met 125	Ser	Cys	Glu	
	cag	aaa	gcc	gaa	gga	cac	ggc		σσα	tet	t ø ø	cag		agt	ttc	gat	432
			Ala														102
		130			,		135		,		•	140					
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		Gln	Ile	Phe	Leu		Phe	Asp	Ser	Glu		Arg	Met	Trp	Thr		
	145		•			150					155	<b>4</b> e				160	500
			cct Pro														528
	val	піS	LI.0	ory	165	wig	LyS	met	LYS	170	LyS	irp	GIU	ASII	175	LyS	
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		_	Thr	_									-	_			

185 190 180 624 tgg ctt gag gac ttc ttg atg ggc atg gac agc acc ctg gag cca agt Trp Leu Glu Asp Phe Leu Met Gly Met Asp Ser Thr Leu Glu Pro Ser 200 679 gca gga ggc aca gtc tgacccaaag ccatggccac caccctcagt ccctgcagcc Ala Gly Gly Thr Val 720 tcctcctcat cctccctgc ttcatcctcc ctggcatctg a <210> 2 <211> 213 <212> PRT <213> Homo sapiens <400> 2 Met Ala Ala Ala Ala Ser Pro Ala Phe Leu Leu Arg Leu Pro Leu Leu 10 5 Leu Leu Ser Ser Trp Cys Arg Thr Gly Leu Ala Asp Pro His Ser 25 Leu Cys Tyr Asp Ile Thr Val Ile Pro Lys Phe Arg Pro Gly Pro Arg 45 40 Trp Cys Ala Val Gln Gly Gln Val Asp Glu Lys Thr Phe Leu His Tyr 60 Asp Cys Gly Ser Lys Thr Val Thr Pro Val Ser Pro Leu Gly Lys Lys 75 Leu Asn Val Thr Thr Ala Trp Lys Ala Gln Asn Pro Val Leu Arg Glu 85 Val Val Asp Ile Leu Thr Glu Gln Leu Leu Asp Ile Gln Leu Glu Asn 105 Tyr Ile Pro Lys Glu Pro Leu Thr Leu Gln Ala Arg Met Ser Cys Glu 120 125 Gln Lys Ala Glu Gly His Gly Ser Gly Ser Trp Gln Leu Ser Phe Asp 135 140 Gly Gln Ile Phe Leu Leu Phe Asp Ser Glu Asn Arg Met Trp Thr Thr 150 Val His Pro Gly Ala Arg Lys Met Lys Glu Lys Trp Glu Asn Asp Lys 170 165 Asp Met Thr Met Ser Phe His Tyr Ile Ser Met Gly Asp Cys Thr Gly 185 190 Trp Leu Glu Asp Phe Leu Met Gly Met Asp Ser Thr Leu Glu Pro Ser 200 Ala Gly Gly Thr Val 210 <210> 3 <211> 29 <212> DNA <213> primer <400> 3 29 cggaattcat ggcagcggcc gccagcccc

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	tcca ggcacagtct ga	742
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